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      Lee N., Kirkness E.F., Weinstock K.G., Gocayne J.D., White O., Sutton G.,
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      Blake J.A., Brandon R.C., Chiu M.-W., Clayton R.A., Cline R.T.,
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      Fritchman J.L., Geoghagen N.S.M., Glodek A., Gnehm C.L., Hanna M.C.,
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     Hedblom E., HinkleJr P.S., Kelley J.M., Klimek K.M., Kelley J.C.,
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      Scott J.L., Saudek D.M., Shirley R., Small K.V., Spriggs T.A.,
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     Utterback T.R., Weidman J.F., Li Y., Bednarik D.P., Cao L., Cepeda M.A., Coleman T.A., Collins E.-J., Dimke D., Feng P., Ferrie A., Fischer C., Hastings G.A., He W.-W., Hu J.-S., Greene J.M., Gruber J., Hudson P.,
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      Kim A., Kozak D.L., Kunsch C., Ji H., Li H., Meissner P.S., Olsen H.,
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      Raymond I., Wei Y.-F., Wing J., Xu C., Yu G.-L., Ruben S.M., Dillon P.J.,
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      "Initial Assessment of Human Gene Diversity and Expression Patterns Based
RT
      Upon 83 Million Basepairs of cDNA Sequence";
RT
      Nature 377:3-174(1995).
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XX
CC
      Contact: Kerlavage, AR
      Bioinformatics
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      The Institute for Genomic Research
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      9712 Medical Center Drive, Rockville, MD 20850 USA
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      Tel: 3018699056
CC
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      Fax: 3018699423
      Email: arkerlav@tigr.org
CC
      For clone availability, additional sequence and expression
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